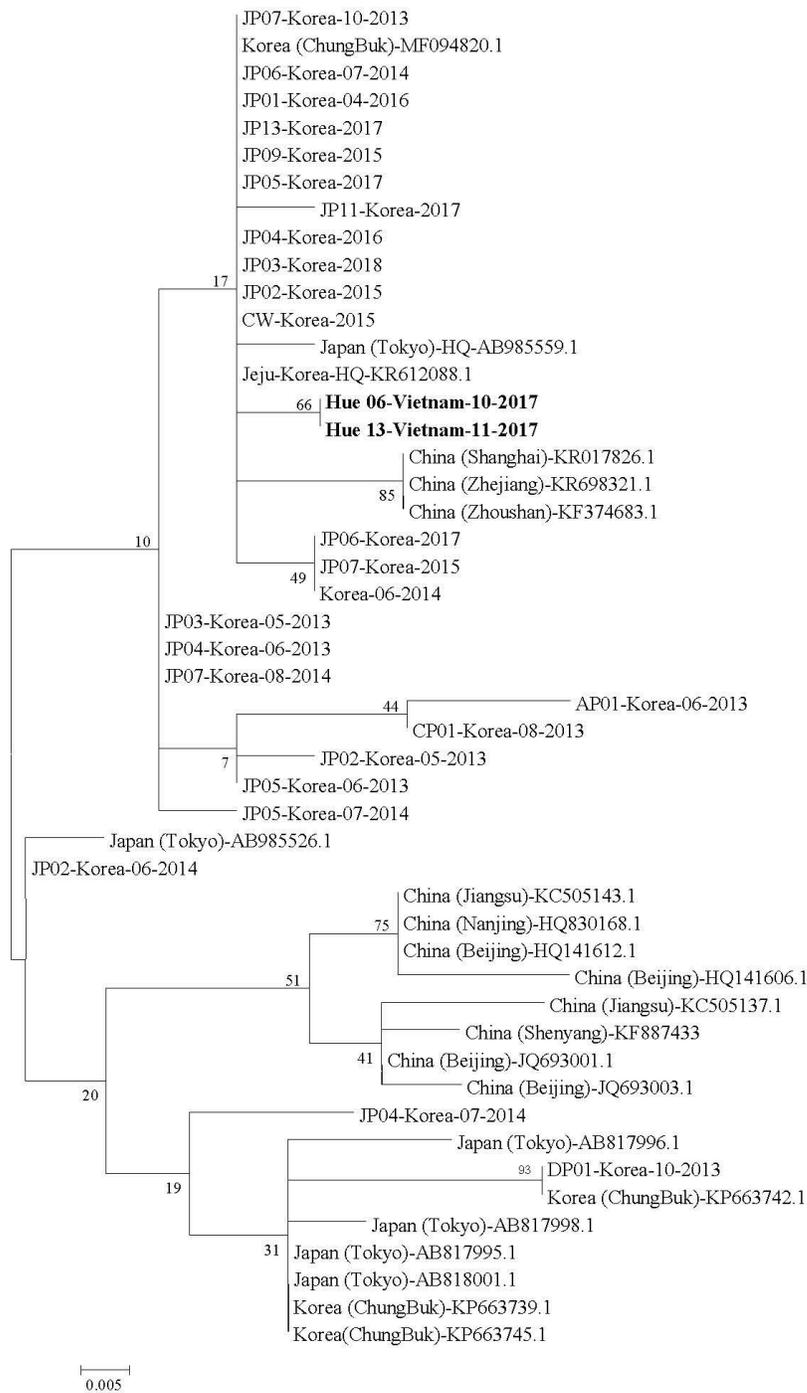


# Endemic Severe Fever with Thrombocytopenia Syndrome, Vietnam

## Appendix

**Appendix Table.** Detection of severe fever with thrombocytopenia syndrome virus antibody, Vietnam

No	Patient	IgM-capture ELISA, optical density	IgG-capture ELISA, optical density
1	Hue 06-Vietnam-10-2017	+3.48	-1.69
2	Hue 13-Vietnam-11-2017	+/-1.05	-1.35
3	Positive control	1.73	2.60
4	Negative control	0.23	0.14



**Appendix Figure.** Phylogenetic tree constructed based on partial small (S) segment sequences of severe fever with thrombocytopenia syndrome virus. The tree was constructed using the maximum-likelihood method with MEGA 6. The partial S sequences from stored serum collected in Vietnam in 2017 and analyzed in this study are shown in bold (Hue-06-Vietnam-10–2017 and Hue-13-Vietnam-11–2017). The partial S sequence data of the viruses identified in China, South Korea, and Japan were obtained from NCBI/BLAST (<http://blast.ncbi.nlm.nih.gov/Blast.cgi>).